Some comments on the paper

"Filter design for the detection of compact sources based on the Neyman-Pearson detector" by M. López-Caniego et. al (2005, MNRAS 359, 993)

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Received; accepted

Abstract. In this note we stress the necessity of a careful check of both the theoretical arguments and the numerical experiments used by López-Caniego et al. (2005) to support the superior performances of the *biparametric scale adaptive filter* (BSAF) with respect to the classic *matched filter* (MF) in the detection of sources on a random Gaussian background.

Key words. Methods: data analysis - Methods: statistical

1. Introduction

A popular tool for the detection of sources embedded in a noise background is the $matched\ filter\ (MF)$. This is because its optimal properties. In particular, in the case of Gaussian noise, it can be shown that through MF it is possible to construct $uniformly\ most\ powerful\ tests$, i.e., tests that for a given $probability\ of\ false\ alarm\ P_{FA}$ (the probability of a false detection) provide the best $probability\ of\ detection\ P_D$ (the probability of a correct detection). This is guaranteed by the Neyman-Pearson (NM) theorem. For the sake of clarity and completeness, some arguments are presented in the following.

Let's suppose that $\boldsymbol{x} = [x(0), x(1), \dots, x(N-1)]^T$ is an observed one-dimensional discrete signal. In its simplest form, the detection problem consists in deciding whether \boldsymbol{x} is constituted by a pure zero-mean Gaussian noise \boldsymbol{n} with covariance matrix \boldsymbol{C} (hypothesis H_0) or it contains also the contribution of a deterministic component \boldsymbol{s} (hypothesis H_1). Both \boldsymbol{C} and \boldsymbol{s} are supposed known. In other words, the source detection problem is equivalent to a decision problem where one has to distinguish between the two hypotheses:

$$\begin{cases}
H_0: & \boldsymbol{x} = \boldsymbol{n}; \\
H_1: & \boldsymbol{x} = \boldsymbol{n} + \boldsymbol{s}.
\end{cases}$$
(1)

Under H_0 the probability of x is given by

$$p(\boldsymbol{x}|H_0) = p_G(\boldsymbol{n}) = p_G(\boldsymbol{x})$$

$$\equiv \frac{1}{(2\pi)^{N/2} \det^{1/2}(\boldsymbol{C})} \exp\left[-\frac{1}{2}\boldsymbol{x}^T \boldsymbol{C}^{-1} \boldsymbol{x}\right], \quad (2)$$

whereas under H_1 it is

$$p(\boldsymbol{x}|H_1) = p_G(\boldsymbol{n}) = p_G(\boldsymbol{x} - \boldsymbol{s})$$

$$\equiv \frac{1}{(2\pi)^{N/2} \det^{1/2}(\boldsymbol{C})} \exp\left[-\frac{1}{2}(\boldsymbol{x} - \boldsymbol{s})^T \boldsymbol{C}^{-1}(\boldsymbol{x} - \boldsymbol{s})\right].$$
(3)

The choice between H_0 and H_1 requires the definition of the so called *critical* or *acceptance region* R_* . This is set of values in \mathbb{R}^N that map into the decision H_1 or

$$R_* = \{ \boldsymbol{x} : \text{decide } H_1 \text{ or reject } H_0 \}. \tag{4}$$

Once R_* has been fixed, the quantities P_D and P_{FA} are given by

$$P_{\rm D} = \int_{R} p(\boldsymbol{x}|H_1)d\boldsymbol{x},\tag{5}$$

and

$$P_{\text{FA}} = \int_{R_*} p(\boldsymbol{x}|H_0) d\boldsymbol{x}. \tag{6}$$

The Neyman-Pearson (NP) theorem tells us how to choose R_* in such a way to maximize P_D when we are given the

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likelihood functions $p(\boldsymbol{x}|H_0)$ and $p(\boldsymbol{x}|H_1)$, and a value α for P_{FA} .

Neyman-Pearson theorem: to maximize P_D for a given $P_{FA} = \alpha$, decide H_1 if the likelihood ratio

$$L(\boldsymbol{x}) = \frac{p(\boldsymbol{x}|H_1)}{p(\boldsymbol{x}|H_0)} > \gamma, \tag{7}$$

where the threshold γ is found from

$$P_{\text{FA}} = \int_{\{\boldsymbol{x}: L(\boldsymbol{x}) > \gamma\}} p(\boldsymbol{x}|H_0) d\boldsymbol{x} = \alpha.$$
 (8)

By inserting Eqs.(5)-(6) in Eq.(7), it is possible to see that H_1 has to be chosen when for the statistic T(x) (called *NP detector*) it is

$$T(\boldsymbol{x}) = \boldsymbol{x}^T \boldsymbol{C}^{-1} \boldsymbol{s} > \gamma', \tag{9}$$

with γ' such as

$$P_{\text{FA}} = Q\left(\frac{\gamma'}{\left\lceil \boldsymbol{s}^T \boldsymbol{C}^{-1} \boldsymbol{s} \right\rceil^{1/2}}\right) = \alpha. \tag{10}$$

Here, $Q(x) = 1 - \Phi(x)$ with $\Phi(x)$ the cumulative distribution function of the standard Gaussian distribution. Equation (10) is due to the fact that, being given by a linear combination of random Gaussian quantities, T(x) is a random Gaussian quantity too. For the same reason it happens that

$$P_{\rm D} = Q \left(Q^{-1} \left(P_{\rm FA} \right) - \sqrt{\frac{\epsilon}{\sigma^2}} \right), \tag{11}$$

with $\epsilon = s^T s$ the signal energy and σ the standard deviation of the noise n. In Eq.(9), the quantity $C^{-1}s$ provides the classical MF. Therefore, this filter can be expected to have optimal properties. An important point to stress is that these properties do not degrade very much when the amplitude and/or the position of the source are unknown and have to be estimated from the data themselves (Kay 1998).

In López-Caniego et al. (2005) (hereafter *LHBS*) an alternative approach is proposed that is claimed to provide filters with superior performances with respect to MF. Authors work in the context of one-dimensional continuous signal and under the hypothesis of sources with symmetric profile. Their idea consists in exploiting the joint distribution $p(\nu_n, \kappa_n)$ of the normalized maxima ν_n and of the corresponding normalized curvature κ_n that, for a continuous Gaussian random noise, is given by the Rice's formula

$$p_R(\nu_n, \kappa_n) = \frac{\kappa_n}{\sqrt{2\pi(1-\rho^2)}} \exp\left[-\frac{\nu_n^2 + \kappa_n^2 - 2\rho\nu_n\kappa_n}{2(1-\rho^2)}\right].$$
(12)

Here, $\nu_n \equiv \xi/\sigma_0 \in (-\infty, +\infty)$, $\kappa_n \equiv -\xi''/\sigma_2 \in (0, +\infty)$ with ξ the value of the maxima of the signal and ξ'' the

corresponding second derivative, whereas σ_i^2 and P(q) are the moment of order 2i

$$\sigma_i^2 \equiv 2 \int_0^\infty q^{2i} P(q) dq, \tag{13}$$

and the power spectrum of the noise process, respectively. Of course, for the first normalized derivative of the maxima it is $\delta_n = \xi'/\sigma_1 = 0$.

If a source is embedded in the background and ν , δ and κ are observed quantities, the NP approach provides the following decision problem

$$\begin{cases} H_0: & \nu = \nu_n, \quad \delta = \delta_n = 0, \quad \kappa = \kappa_n; \\ H_1: & \nu = \widetilde{\nu}_n + \widetilde{\nu}_s, \quad \delta = \widetilde{\delta}_n + \widetilde{\delta}_s = 0, \quad \kappa = \widetilde{\kappa}_n + \widetilde{\kappa}_s, \end{cases}$$
(14)

where $\widetilde{\nu}_n$, $\widetilde{\nu}_s$ are the values of the noise process and of the source in correspondence to the position of the observed maximum, whereas $\widetilde{\delta}_n$, $\widetilde{\delta}_s$ and $\widetilde{\kappa}_n$, $\widetilde{\kappa}_s$ are the corresponding normalized first derivatives and curvatures, respectively. In other words, under H_0 it is $p(\nu,0,\kappa|H_0)=p_R(\nu_n,\kappa_n)$, whereas under H_1 it is $p(\nu,0,\kappa|H_1)=p_f(\widetilde{\nu}_n,\widetilde{\delta}_n,\widetilde{\kappa}_n)$. A point to stress is that the likelihood $p_f(\widetilde{\nu}_n,\widetilde{\delta}_n,\widetilde{\kappa}_n)$ is not given by the Rice's formula. This is because, in general, there is no guarantee that the position of the observed maximum coincides with the position of the peak of the source and/or with the position of one of the peaks in the noise process.

2. Some comments on the LHBS approach

The *LHBS* approach deserves some comments. Among these, three are of particular relevance. The first one is that, instead of $p_f(\tilde{\nu}_n, \tilde{\delta}_n, \tilde{\kappa}_n)$, under H_1 authors use a likelihood in the form

$$p(\nu, \kappa | H_1) = \frac{\kappa}{\sqrt{2\pi(1 - \rho^2)}} \times \exp\left[-\frac{(\nu^2 - \nu_s)^2 + (\kappa - \kappa_s)^2 - 2\rho(\nu - \nu_s)(\kappa - \kappa_s)}{2(1 - \rho^2)}\right],$$
(15)

where $\nu \in (-\infty, +\infty)$ and $\kappa \in (0, +\infty)$. Actually, it is not clear from where this equation comes out. In fact, even in the unrealistic hypothesis that the position of the peak of the source has to coincide with the position of a maximum of the noise process, similarly to Eq.(3) it should be

$$p(\nu, \kappa | H_1) = \frac{\kappa - \kappa_s}{\sqrt{2\pi (1 - \rho^2)}}$$

$$\times \exp\left[-\frac{(\nu^2 - \nu_s)^2 + (\kappa - \kappa_s)^2 - 2\rho(\nu - \nu_s)(\kappa - \kappa_s)}{2(1 - \rho^2)}\right],$$
(16)

with $\nu \in (-\infty, +\infty)$ and $\kappa \in (\kappa_s, +\infty)$. LHBS justify the use of the likelihood (15) only with a reference to Barreiro et al. (2003). However, also in that work no details are provided.

The second comment regards the fact that LHBS do not work on the original signal x, but on a filtered version of it. The reason is that this operation modifies R_* (see Fig. 5 in Barreiro et al. 2003). Hence, LHBS try to define a filter for which, once fixed $P_{\rm FA}$, R_* is such as to maximize the corresponding $P_{\rm D}$. This is accomplished by parametrizing some well known filters (e.g., MH and the Mexican hat wavelet) and by computing the values of the parameters through an optimization procedure. For example, the biparametric scale adaptive filter (BSAF), claimed by LHBS to outperform MH, is a modified version of MF where two coefficients are assumed to be free parameters. However, authors provide only an heuristic explanation of the reason why such an approach should be expected to work. No rigorous argument is given. Probably, the conjecture is that their procedure provides a two-steps (parameter optimization + filtering) approximation of the solution of the decision problem based on the likelihood ratio

$$L(\boldsymbol{x}, \nu, \kappa) = \frac{p(\boldsymbol{x}, \nu, \kappa | H_1)}{p(\boldsymbol{x}, \nu, \kappa | H_0)} > \gamma, \tag{17}$$

that (at least in principle) should better exploit the available a priori information. The main problem to work directly with such likelihood ratio is in the difficulty to obtain the joint distributions $p(\boldsymbol{x}, \nu, \kappa | H_0)$ and $p(\boldsymbol{x}, \nu, \kappa | H_1)$ that can be quite complex. In any case, the usefulness of the proposed procedure is not obvious and should be proved through rigorous theoretical arguments. The conclusions drawn only on the basis of numerical experiments lack generality and can be potentially misleading. This is especially true for the discrete case where the quantities ν and κ have to be computed by using the data \boldsymbol{x} .

The last comment regards the procedure followed by authors in the numerical experiments. In particular, it is questionable that they investigate the performance of their detection test only on the sources that, after the filtering, show a peak placed at the position of their true maximum. This is certainly an unrealistic condition. Moreover, in this way there is the risk to favour the selection of the sources that are located in correspondence to a maximum of the noise, i.e., those whose likelihood $p(\nu, \kappa|H_1)$ is given by Eq.(16). In the discrete case, it easy to realize that this is very probable when, for example, a source is superimposed on a white noise background and no filtering operation is carried out. On the other side, BSAF is a symmetric filter that works principally on spatial scales smaller than a pixels. Hence, the necessity to check for the above mentioned risk. The correct procedure requires that the detection test be applied also to the sources that after the filtering show a maximum at a position different from the true one.

3. Conclusions

From the discussion presented above, it is evident the necessity of a careful check of both the theoretical arguments and the numerical experiments used by *LHBS* before BSAF can be claimed to be superior to the classic MH in source detection problems.

As last comment, it is useful to stress that the BSAF corresponding to a specific problem can be obtained only through a very involved procedure based on stringent *a priori* assumptions (e.g., Gaussianity of the background, symmetry of the source profile). All this makes the procedure proposed by *LHBS* quite rigid. In this respect, MF certainly represents a much more flexible solution.

References

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López-Caniego, M., Herranz, D., Barreiro, R.B., & Sanz, J.L. 2005, MNRAS, 359, 993 (*LHBS*)